PCT/FR00/00144

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į	Names	similarities	Repeats	ORFs
- 7582	Recons. RNA		yes	538
- 37879	RG083M05 [7]	%96	yes	538
14079	BAC378 [14]	88%	yes	OL OL
27999	Q11M15 [21]		yes	413 and 305
94627		88%	2	٤

REPLACEMENT SHEET (RULE 26)

10 20 30 40 50 1234567890 1234567890 1234567890 1234567890	
OCTAGAACGT ATTCTGGAGA ATTGGGACCA ATGTGACACT CAGACGCTAA PRTYSGELGPM.HSDAK I.ERILENWDQCDTQTLR .NVFWRIGTNVTLRR	50
CAAACAAACG ATTTATATTC TICIOCAGTA CCOCCTGOCC ACAATATCCT KETIYILLQYRLA TISS KKRFIFFCSTAWPQYP ERNDLYS SAVPPGHNIL	100
CTICAAGGGA GAGAAACCTG OCTTCCTGAG GGAAGTATAA ATTATAACAT SRERNLAS.GKYKL.H LQGRETWLPEGSINYNI FKGEKPGFLREV.IITS	150
CATCTTACAG CTAGACCTCT TCTGTAGAAA GGAGGGCAAA TGGAGTGAAG H L T A R P L L . K G G Q M E . S I L Q L D L F C R K E G K W S E V S Y S . T S S V E R R A N G V K	200
TOCCATATET GCAAACITTC TTITCATTAA GAGACAACIC ACAATTATGT A I C A N F L F I K R Q L T I M . P Y V Q T F F S L R D N S Q L C C H M C K L S F H . E T T H N Y V	250
AAAAAGIGIG GITTIATGOOC TACAGGAAGC CCICAGAGIC CACCICOCTA K V W F M P Y R K P S E S T S L K K C G L C P T G S P Q S P P P Y K S V V Y A L Q E A L R V H L P T	300
COCCAGGIC COCTOCOGA CICCITOCIC ANCIAATAAG GACCCCCTT PQRPLPDSFLNGPPF PSVPSPTPSSTNKDPPL PASPPRLLPQLIRTPL	350
TRACOCRARC GOTOCRARAG CREATRGREA ARGOGGTRARA CRATCARCCA N P N G P K G D R Q R G K Q . T K T Q T V Q K E I D K G V N N E P . P K R S K R R . T K G . T M N Q	400
AACAGIOCCA ATTATICCCCC ATTATICCCCC CICCAACCAG TCACACGAGG E C Q Y S P I M P P P S S E R R K S A N I P R L C P L Q A V R G G R V P I F P D Y A P S K Q . E E E	450
AGAATTOOCC CCAGCCAGAG TOCCTGTACC TITTTCCTCC TCAGACTTAA RIRPSQSACTFFSLRLK EFGPARVPVPFSLSDLK NSAQPECLYLFLSQT.	500

REPLACEMENT SHEET (RULE 26)

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FIG 2 (continued)

10 20 30 40 50 1234567890 1234567890 1234567890 1234567890	
ACCAAATTIAA AATAGACCTA OGTAAATTICT CAGATAACCC TGACGGCTAT A N N R P R . I L R . P . R L Y Q I K I D L G K F S D N P D G Y S K L K . T . V N S Q I T L T A I	550
ATTGATGTTT TACAAOGGIT AGGACAATCC TITGATCTGA CATGGAGAGA C F T R V R T I L S D M E R I D V L Q G L G Q S F D L T W R D L M F Y K G D N P L I . H G E I	600
TATAATGITA CIACTAAATC AGACACTAAC CCCAAATGAG AGAAGTOCOG Y N V T T K S D T N P K . E K C R I M L L L N Q T L T P N E R S A A . C Y Y . I R H . P Q M R E V P	650
CIGTAACTOC ACCOCCACAG TITICOCCATC TITICGITATCT CAGICAGOCC C N C S P R V W R S L V S Q S G Q V T A A R E F G D L W Y L S Q A L . L Q P E S L A I F G I S V R P	700
AACAATAGGA TGACAACAGA GGAAAGAACA ACTOOCACAG GOCAGCAGGC Q . D D N R G K N N S H R P A G N N R M T T E E R T T P T G Q Q A T I G . Q Q R K E Q L P Q A S R Q	750
ACTICCCACT CIPACACCIC ATTOCCACAC ACASTCACAA CATGCACATT S S Q C R P S L G H R I R T W R L V P S V D P H W D T E S E H G D W F P V . T L I G T Q N Q N M E I	800
GCTGCCACAA ACATTTGCTA ACTTGCGTGC TAGAAGGACT GAGGAAAACT V P Q T F A N L R A R R T E E N . C H K H L L T C V L E G L R K T G A T N I C . L A C . K D . G K L	850
ACCAACAACC CTATGAATTA CICAATGATG TOCACTATAA CACAGGCAAA EEAYELLNDVHYNTGK RKKPMNYSMMSTITQGK GRSL.ITQ.CPL.HRER	900
GRAGAAAT CTEACTOCIT TICTGCACAG ACTAAGGGAG GCATTICAGGA GRKS Y C F S G O T K G G I E E E E N L T A F L D R L R E A L R K K K I L L F W T D . G R H . G	950
ACCATACCTC CCTGTCACCT CACTCTATTG AACCCCAACT AATCTTAAAG A Y L P V T . L Y . R P T N L K G H T S L S P D S I E G Q L I L K S I P P C H L T L L K A N . S . R	1000

20 30 40 10 1234567890 1234567890 1234567890 1234567890 1234567890 CATAAGITTA TCACTCAGIC AGCTOCAGAC ATTAGAAAAA ACTTCAAAAG . VY HSV SCRH . KK LQK DKFI TQS AAD IRKN FKS ISL SLSO LOT LEK TSKV TCTGCCTTAG GCCCGGAGCA GAACTIVAGAA ACCCUATTEA ACTTGGCATC SALG PEQ NLE TLFN LAS LP. ARSRT.KPYLTWHP CLR PGA ELRN PI. CTCAGTTTTT TATAATAGAG ATCAGGAGGA GCAGGGGAAA CGGGACAAAC SVFYNRD QEE QAK RDKR Q F F I I E I R R S R R N G T N LSFL . R SGG AGET GQT GCCATAAAAA AAAAAGCCCG GCTCCACTAC TTTAGTCATG GCCCTCAGGC 1200 DKKKRGGPLL.SWPSGGIKKKGGVHYFSHGPQAG.KKKGGSTTLVMALRQ AAGCAGACIT TGGAGGCTCT GCAAAAGGGA AAAGCTGGGC AAATCAAATG KQTL EAL QKG KAGQ IKC SRL WRLC KRE KLG KSNA ADF GGS AKGK SWA NQ M CCTAATAGGG CTGGCTTCCA GTGCGGTCTA CAAGGACACT TTAAAAAAAGA L I G L A S S A V Y K D T L K K I . . G W L P V R S T R T L . K R PNRAGFQ CGL QGHF KKD TTATOCAAGT ACAAATAAGC CGCCCCTTG TCCATGCCCC TTACGTCAAG IQV EIS RPLV HAP YVK LSK. K.AAPL SMPL TSR Y P S R N K P P P C P C P L R Q G GGAATCACTG GAAGGCCCAC TGCCCCAGGG GATGAAGATA CTCTGAGTCA GITG RPT APG DEDT LSQ ESLEGPL PQG M K I L . V R NHWKAH CPRG . RYSES CANGOUNTIN NOUNCATIONT CONCUNCYG CACTGAGOGT GCCCCGGGGGG SH. PDD PAAG LRV PGA EAIN QMI QQQ D. GC PGR AGOSCOAGO: CATGOCATCA COCTCACAGA COCCOGGIA TGTTTGACCA R Q P M P S P S Q S P G Y V . P S A S P C H H P H R A P G M F D H APA HAIT LTE PRV CLTI

REPLACEMENT SHEET (RULE 26)

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
TTGAGAGCCA					1511

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